## SEQUENCE LISTING

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<110> MILLENNIUM PHARMACEUTICALS, INC.
<120> NOVEL GENES ENCODING PROTEINS HAVING DIAGNOSTIC,
      PREVENTIVE, THERAPEUTIC, AND OTHER USES
<130> 210147.0034/12WO
<140> Not Yet Assigned
<141> 2000-06-30
<150> US 09/345,293
<151> 1999-06-30
<150> US 09/345,680
<151> 1999-06-30
<160> 35
<170> PatentIn Ver. 2.1
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gagtctggcc tatctcagag ttgccagcct ttccctgtct accagaccat ggaggtccag 540
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Leu Val Leu Gly Leu Ser Leu Cys Val His Leu Leu Met Phe Pro Ala
         35
                             40
Ala Val Ser Gly Gly Arg Asp Cys His Ile Leu Glu His Gly His Ile
Tyr Leu His Arg Cys Thr Glu Pro Asp Met Leu Pro Phe Glu Ser Gln
 65
                     70
                                          75
His Ser Val Thr Thr Leu Ser Ile Ala Pro Glu Arg Gly Pro Glu Pro
Thr Ala Leu His Ser Cys Leu His Lys Asn Arg Asp Gly Leu Gly Arg
            100
                                105
                                                     110
Ile Met Lys Arg Lys Ile Trp Leu Ala Leu Gly Leu Gly His His Ile
                            120
Gly Ser Ala Ala Tyr Cys Val Ser Gly Ser Arg Ala Pro Trp Glu Ala
Ser Cys Gly Cys His Asp Ser Ser His Pro Trp Ser Ser His Ser Met
145
                    150
Glu Ser Gly Leu Ser Gln Ser Cys Gln Pro Phe Pro Val Tyr Gln Thr
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Arg Thr Gly Glu
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<213> Homo sapiens

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Ala Val Ser Gly Gly 50

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<211> 143

<212> PRT

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Thr Glu Pro Asp Met Leu Pro Phe Glu Ser Gln His Ser Val Thr Thr 20 30

Leu Ser Ile Ala Pro Glu Arg Gly Pro Glu Pro Thr Ala Leu His Ser

Cys Leu His Lys Asn Arg Asp Gly Leu Gly Arg Ile Met Lys Arg Lys 50

Ile Trp Leu Ala Leu Gly Leu Gly His His Ile Gly Ser Ala Ala Tyr

Cys Val Ser Gly Ser Arg Ala Pro Trp Glu Ala Ser Cys Gly Cys His 90 95

Asp Ser Ser His Pro Trp Ser Ser His Ser Met Glu Ser Gly Leu Ser 105 110

Gln Ser Cys Gln Pro Phe Pro Val Tyr Gln Thr Met Glu Val Gln Pro 120

Tyr Met Ser Cys Gly Pro Thr Asp Gln Leu Val Arg Thr Gly Glu 135

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<211> 58

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Phe Pro Val Tyr Gln Thr Met Glu Val Gln Pro Tyr Met Ser Cys Gly Pro Thr Asp Gln Leu Val Arg Thr Gly Glu <210> 7 <211> 21 <212> PRT <213> Homo sapiens Ile Trp Leu Ala Leu Gly Leu Gly His His Ile Gly Ser Ala Ala Tyr Cys Val Ser Gly Ser 20 <210> 8 <211> 64 <212> PRT <213> Homo sapiens <400> 8 Arg Asp Cys His Ile Leu Glu His Gly His Ile Tyr Leu His Arg Cys 10 Thr Glu Pro Asp Met Leu Pro Phe Glu Ser Gln His Ser Val Thr Thr 20 30 Leu Ser Ile Ala Pro Glu Arg Gly Pro Glu Pro Thr Ala Leu His Ser Cys Leu His Lys Asn Arg Asp Gly Leu Gly Arg Ile Met Lys Arg Lys 55 60 <210> 9 <211> 22 <212> PRT <213> Homo sapiens <400> 9 Leu Phe Phe Pro Thr Arg Leu Gln His Pro Leu Arg Glu Gly Asp Gln 10 Asn Arg Gly Pro Ser Ser 20

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<211> 67
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His Arg Cys Thr Glu Pro Asp Met Leu Pro Phe Glu Ser Gln His Ser
             20
                                                      30
Val Thr Thr Leu Ser Ile Ala Pro Glu Arg Gly Pro Glu Pro Thr Ala
Leu His Ser Cys Leu His Lys Asn Arg Asp Gly Leu Gly Arg Ile Met
     50
                         55
Lys Arg Lys
65
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acaaggaget ggetttggge taggetgete ettgeetatg attggggaag gttaaacece 180
tacagggctt atgtatgtgg aaactgttgg aacactgatt aaatgggatg gacttcactt 240
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taateettta qqqaqqeaqa aaaqqeeaqa atqcaaaqee atetttteat tacactaqqq 360
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agtgageetg geeaggacag taacetgtgg gettgtgatg acattattte taatagggaa 480
tgggaaagga tgttagcttc tcaggtttta aagtgtcctg gaggagaaga gaaaggacga 540
catqaqaaqq aqacaatqaa gaaqatqqqt qaqqqqqaqa tagtqtaaqa ccctqaqaat 600
qqcataqqqt aaaactqqqa caqaqatact qtqqqaqaac qataqctqca qaqqqacaga 660
gggaggaagg aaggagaaga gagggagata aaaacagttt qqaqaaactc tcacaataca 720
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<213> Homo sapiens

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Trp Ala Phe Ile Trp Gly Gly His Val Ser Pro Thr Trp Asn Ser Glu
20 25 30

Pro Gly Gln Asp Ser Asn Leu Trp Ala Cys Asp Asp Ile Ile Ser Asn 35 40 45

Arg Glu Trp Glu Arg Met Leu Ala Ser Gln Val Leu Lys Cys Pro Gly 50 55 60

Gly Glu Glu Lys Gly Arg His Glu Lys Glu Thr Met Lys Lys Met Gly 65 70 75 80

Glu Gly Glu Ile Val

<210> 24

<211> 23

<212> PRT

<213> Homo sapiens

<400> 24

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<211> 62

<212> PRT

<213> Homo sapiens

<400> 25

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Trp Ala Cys Asp Asp Ile Ile Ser Asn Arg Glu Trp Glu Arg Met Leu 20 25 30

Ala Ser Gln Val Leu Lys Cys Pro Gly Gly Glu Glu Lys Gly Arg His
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Glu Lys Glu Thr Met Lys Lys Met Gly Glu Gly Glu Ile Val
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<210> 27

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cettgtgccc tcccagggct tcagctggtg gcgctggctg cttgactttc tggaaatgga 240
tgccatgcaa ggttgcattt tgaccagggg gaaggaggtc attcttgagg gtggctggat 300
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Ile Asp Cys Glu Ser Ala Pro Ala Pro Glu Thr Arg Gln Gly Cys Leu  $35 \hspace{1cm} 40 \hspace{1cm} 45$ 

Glu Ala Pro Pro Pro Ser Glu Thr Asp Ser Gln Gln Ser His Thr Val
50 55 60

Trp Ser Ala Lys Cys 65